



## Short communication

## Actin-independent trafficking of cochlear connexin 26 to non-lipid raft gap junction plaques

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## ABSTRACT

Hereditary hearing loss affects about 1 per 1000 children. Mutations in GJB2, which encodes the connexin 26 protein (Cx26) involved in cochlear homeostasis, are found in about 50% of patients with autosomal recessive non-syndromic hearing loss. Deciphering the trafficking pathway of cochlear Cx26 *in situ* should represent an advance in understanding the pathogenic significance of many of these mutations. Connexins trafficking and delivery to lipid raft-associated gap junction plaques usually requires successively microtubule and actin networks. Here we show that cochlear Cx26 exhibits an unusual trafficking pathway. We observed that Cx26 assembly occurs in non-lipid raft membrane domains and that junctional plaques are devoid of actin and associated zonula occludens proteins. Using cytoskeleton-disrupting drugs in organotypic culture, we found that cochlear Cx26 gap junction assembly requires microtubules but not actin filaments. Altogether, our data provide an unexpected insight into Cx26 trafficking pathway and gap junction assembly in the cochlea.

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## 1. Introduction

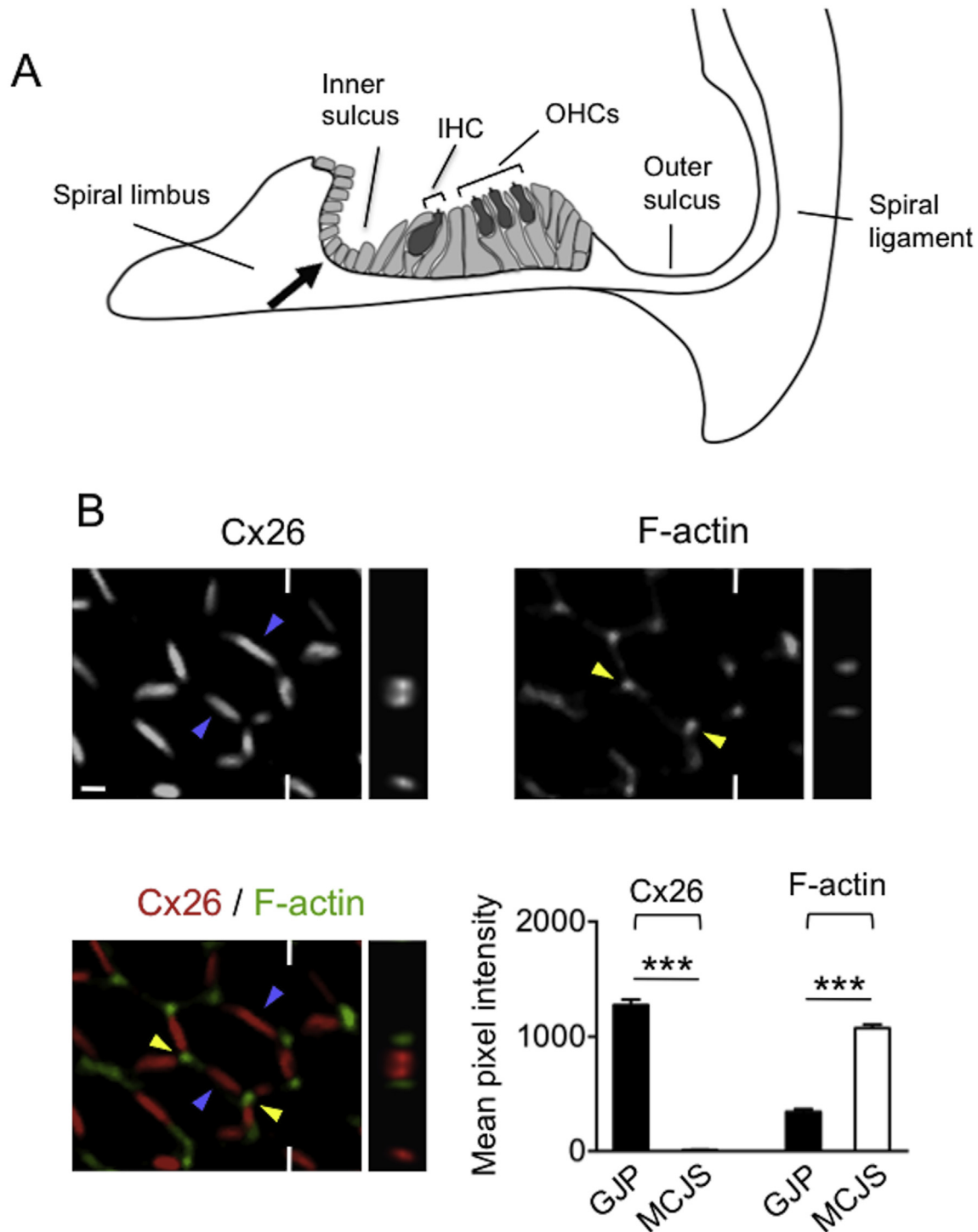
Hearing loss is the most common congenital sensory deficit. About 1–3 in 1000 children are affected at birth or during early childhood by severe hearing loss, which is defined as prelingual deafness, with at least half of all cases attributable to genetic causes (Korver et al., 2017). Despite extraordinary genetic heterogeneity, mutations in one gene, *GJB2*, which encodes the connexin 26 protein (Cx26) and is involved in inner ear homeostasis, are found in up to 50% of patients with autosomal recessive non-syndromic hearing loss (Kelsell et al., 1997). Beside this non-syndromic form of deafness, *GJB2* mutations cause several types of syndromic hearing loss associated with skin diseases with variable prognosis (Avshalumova et al., 2014; Lilly et al., 2016; Xu and Nicholson, 2013). In mammals, sounds are perceived through mechano-sensory hair cells located within the sensory epithelium of the cochlea (i.e. the organ of Corti). Within the organ of Corti, sensory inner and outer hair cells and non-sensory supporting cells are organized in a regular mosaic pattern that extends along the basal-to-apical axis of the cochlear duct. Cx26 gap junction protein, which assembles to form channels between cochlear supporting cells,

allows the rapid removal of K<sup>+</sup> away from the base of sensory hair cells, resulting in the recycling of this ion back to the endolymph to maintain cochlear homeostasis (Kikuchi et al., 2000). However, gap junctions may serve additional roles in the cochlea, such as providing networks for nutrient transfer (Chang et al., 2008; Jagger and Forge, 2015). Cx26 and Cx30 (encoded by the deafness gene *GJB6* (Grifa et al., 1999)) are the two most abundantly expressed gap junction proteins in the cochlea and form heteromeric and heterotypic channels between adjacent supporting cells, from the spiral limbus to the cochlear spiral ligament (Ahmad et al., 2003; Sun et al., 2005) (Fig. 1A). Although the two channel components Cx26 and Cx30 are well characterized, the gap junction plaque (GJP) assembly mechanisms occurring *in situ* remain largely unknown. Of note, however, is the fact that these two proteins are not functionally equivalent, since Cx26 has been shown as the key organizer of the gap junction macromolecular complex (Kamiya et al., 2014). Beside mutations that affect the Cx26 channel function itself, many of the disease-causing mutations in *GJB2* impair the trafficking and delivery of Cx26 to the cell surface, what prevents the formation of gap junctions (Ambrosi et al., 2013; Hoang Dinh et al., 2009; Xu and Nicholson, 2013). Thus, deciphering the trafficking pathway of cochlear Cx26 should represent an advance in understanding the pathogenic significance of these mutations.

Gap junction assembly usually occurs in a “two-step mechanism” which requires successively microtubules and actin

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**Fig. 1. Cx26 GJPs are devoid of F-actin.** (A) Schematic representation of a transverse section through a single cochlear turn. The region of interest (inner sulcus) is indicated by an arrow. (B) Cx26 immunolabelling and F-actin staining of inner sulcus cells of a newborn mouse. F-actin (yellow arrowheads) does not associate with Cx26 (blue arrowheads) but localizes at multicellular junction sites between adjacent GJPs. Mean pixel intensity measurement shows that Cx26 immunolabelling is stronger in the GJP and absent from the multicellular junction sites ( $n = 10$ ). In contrast, the cortical actin network is weaker in the GJP itself, and enriched at the multicellular junction sites ( $n = 10$ ). Statistical significance was determined using Student's *t*-test. Data are presented as mean  $\pm$  SEM. \*\*\* $P < 0.001$ . Scale bar represents 1  $\mu$ m. GJP = gap junction plaque; IHC = inner hair cell; OHCs = outer hair cells; MCJS = multicellular junction site. (For interpretation of the references to colour in this figure legend, the reader is referred to the Web version of this article.)

cytoskeletal components. First, hexameric connexons assembled in the *trans*-Golgi network are trafficked along microtubules to the non-junctional plasma membrane (Koval et al., 1997; Lauf et al., 2002; Musil and Goodenough, 1993). Secondly, hemichannels associate with cortical actin through actin-binding proteins zonula occludens (ZOs) which regulate delivery of connexins from the

periphery to the GJP (Hervé et al., 2014; Thévenin et al., 2013). The membrane lipid environment is critical for gap junction assembly and function (Cascio, 2005; Defamie and Mesnil, 2012). Junctional connexins assemble into lipid raft microdomains whereas non-junctional connexins are present in non-lipid raft fractions (Defamie and Mesnil, 2012; Hunter et al., 2005; Musil and

Goodenough, 1991). Because of the relatively short half-life of connexins (usually 1 – 5 h), the junctional plaque is in a dynamic state, constantly remodeled through both recruitment of newly synthesized connexons to the periphery and endocytosis of older components from the center of the plaque (Gaietta et al., 2002).

Here we show that cochlear Cx26 exhibits an unexpected trafficking pathway towards non-lipid raft gap junction plaques. We observed that junctional Cx26 are devoid of F-actin and associated ZO-1 and ZO-2. Using cytoskeleton-disrupting drugs in organotypic culture, we found that the assembly of Cx26 into GJPs requires microtubules but not actin networks. Therefore, unlike most connexins, the trafficking of cochlear Cx26 towards non-lipid raft GJPs occurs in a kind of “one-step mechanism”, only microtubule-dependent.

## 2. Materials and methods

### 2.1. Animals

Mice of the BALB/c strain were group-housed in the animal facility of the University of Liège under standard conditions with food and water ad libitum and were maintained on a 12-h light/dark cycle. All animals were taken care in accordance with the Declaration of Helsinki and following the guidelines of the Belgian ministry of agriculture in agreement with EC laboratory animal care and use regulation (2010/63/UE, 22 September 2010).

### 2.2. Tissue processing and immunostainings

Cochleae of newborn mice were fixed for 2 h in 4% formaldehyde. Whole-mount cochleae or organotypic explants were incubated overnight at 4°C with primary antibodies directed against connexin 26 (rabbit polyclonal antibody; 1:500; Invitrogen), ZO-1 (rat monoclonal antibody, 1:50; Santa Cruz Biotechnology) and ZO-2 (mouse monoclonal antibody, 1:50; Santa Cruz

Biotechnology). TRITC-conjugated phalloidin (1:500; Sigma-Aldrich) was used as an F-actin marker. Lipid rafts were labelled using FITC-conjugated Cholera Toxin B subunit (1 µg/mL; Sigma-Aldrich). Tissues were then incubated for 1 h with either Rhodamine Red X- or FITC-conjugated goat anti-mouse, anti-rabbit or anti-rat IgGs secondary antibodies (Jackson ImmunoResearch Laboratories).

### 2.3. In vitro organotypic assay

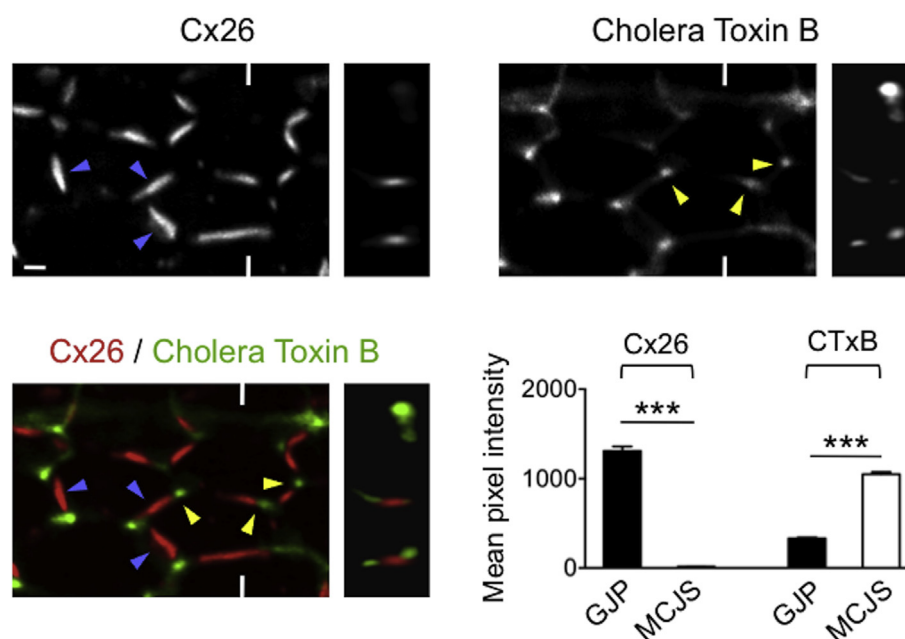
Organs of Corti were isolated from newborn mice and cultured onto Millicell Culture Insert (Millipore) as previously described (Defourny et al., 2015). Organotypic cultures were incubated for 2 h with dimethyl sulfoxide (DMSO, vehicle), nocodazole (100 µM; Sigma-Aldrich) or cytochalasin D (10 µM; Sigma-Aldrich).

### 2.4. Confocal microscopy, image analysis and quantification

Confocal fluorescence images were acquired using the Olympus Fluoview FV1000 confocal system (Olympus Europa GmbH). For comparison between different culture conditions, all preparations were analysed at the same time, using the same acquisition parameters. For each culture condition, 150 inner sulcus cells and 150 GJPs were quantified and measured from three independent experiments and data were plotted. Inner sulcus cells were randomly chosen and GJPs were measured using ImageJ software. Mean pixel intensities in GJPs and in adjacent multicellular junction sites were measured using ImageJ software.

### 2.5. Statistics

All data are presented as mean ± SEM. Data were statistically analysed using two-tailed Student's *t*-test or one-way ANOVA followed by Dunnett's post-test. *P*-values < 0.05 were considered significant (\*\*\*) *P* < 0.001).



**Fig. 2. Cx26 assembles into non-lipid raft GJPs.** Cx26 immunolabelling and Cholera Toxin B staining of inner sulcus cells of a newborn mouse. Lipid rafts (yellow arrowheads) do not associate with Cx26 (blue arrowheads) but localize at multicellular junction sites between adjacent GJPs. Mean pixel intensity measurement shows that Cx26 immunolabelling is stronger in the GJPs and absent from the multicellular junction sites ( $n = 10$ ). In contrast, the lipid rafts are absent from the GJP itself, and enriched at the multicellular junction sites ( $n = 10$ ). Statistical significance was determined using Student's *t*-test. Data are presented as mean ± SEM. \*\*\* $P < 0.001$ . Scale bar represents 1 µm. GJP = gap junction plaque; MCJS = multicellular junction site. (For interpretation of the references to colour in this figure legend, the reader is referred to the Web version of this article.)

## 2.6. Data availability

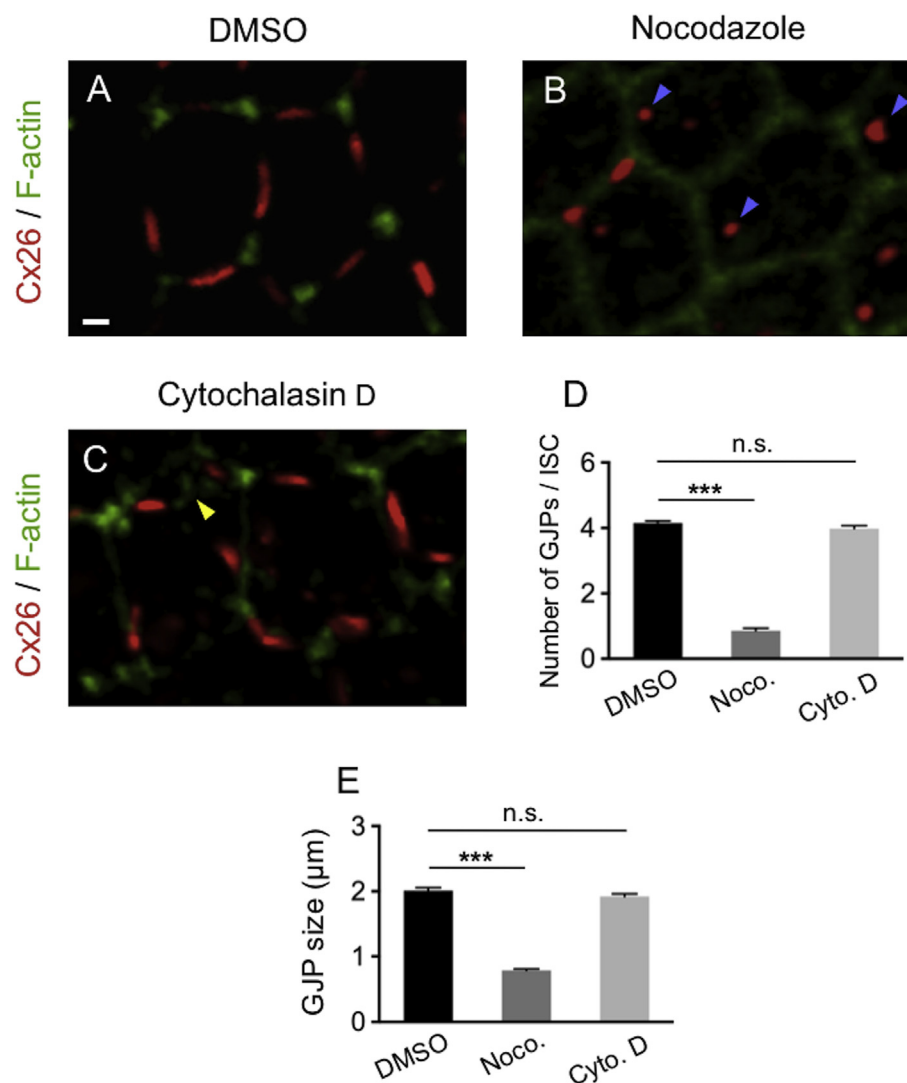
The data that support the findings of this study are available from the corresponding author upon reasonable request.

## 3. Results and discussion

### 3.1. Cx26 gap junctions are devoid of cortical actin and associated ZOs

Once they have reached the plasma membrane, most connexins are linked to the submembrane actin network through ZOs. ZOs contain three PDZ domains and connexins interact with them via a PDZ-binding motif (Hervé et al., 2014; Thévenin et al., 2013). Interestingly, the genomic duplication and overexpression of TJP2/ZO-2 causes a progressive non-syndromic hearing loss in humans (Walsh et al., 2010). We then examined whether cochlear Cx26 gap

junctions associate with cortical actin and ZO proteins in inner sulcus cells of newborn mice. These non-sensory cells, located close to the inner hair cell layer, were previously considered to address the role of Cx26 in the assembly of the gap junction macromolecular complex in the cochlea (Kamiya et al., 2014) (the inner sulcus is indicated by an arrow in Fig. 1A). Surprisingly, we found that F-actin does not associate with Cx26 but rather localizes to multicellular junctions sites between adjacent GJPs (Fig. 1B). Similar distributions were observed in outer sulcus cells, which are non-sensory cells located between the outer hair cell rows and the spiral ligament (Fig. S1A). F-actin and Cx26 therefore display mutually exclusive localizations. Moreover, a co-immunolabelling of Cx26 and ZO-1 or ZO-2 revealed that none of these two proteins associate with junctional Cx26 (Fig. S2). Although rather surprising, these findings are consistent with the fact that murine and human Cx26 do not contain any predicted PDZ-binding motif at their C-terminal extremity (Thévenin et al., 2013; Sheng and Sala, 2001).



**Fig. 3. Cx26 assembly into GJPs does not require actin networks.** (A–C) Organotypic cultures of organ of Corti were treated with DMSO as a control condition, nocodazole or cytochalasin D. (B) Nocodazole treatment strongly disrupts the formation of GJPs and leads to intracellular accumulation of Cx26 (blue arrowheads). In this condition, actin networks extend towards bicellular junctions. (C) The treatment with cytochalasin D does not affect the formation of Cx26 GJPs. The effect of cytochalasin D on actin networks was ascertained by the presence of actin aggregates throughout the cytoplasm (yellow arrowhead). (D) The number of GJPs per inner sulcus cell is significantly reduced by nocodazole but not by cytochalasin D treatment as compared to control condition ( $n = 150$  inner sulcus cells per condition;  $***P < 0.001$ ). (E) The size of GJPs is significantly reduced by nocodazole but not by cytochalasin D treatment as compared to control condition ( $n = 150$  GJPs per condition;  $***P < 0.001$ ). Statistical significance was determined using one-way ANOVA test followed by Dunnett's post-test. Data are presented as mean  $\pm$  SEM. n.s. = not significant. Scale bar in (A) represents  $1 \mu\text{m}$  in (A–C). GJP = gap junction plaque; ISC = inner sulcus cell. (For interpretation of the references to colour in this figure legend, the reader is referred to the Web version of this article.)



The cytoplasmic tail is the most divergent domain with varying sizes among the connexins and is believed to interact with a variety of molecules (Kotini and Mayor, 2015). Since Cx26 is one of the smallest connexins with a short C-terminal tail, this could explain why Cx26 cannot associate with F-actin and/or actin-binding proteins.

### 3.2. Cx26 assembles into non-lipid raft gap junction plaques

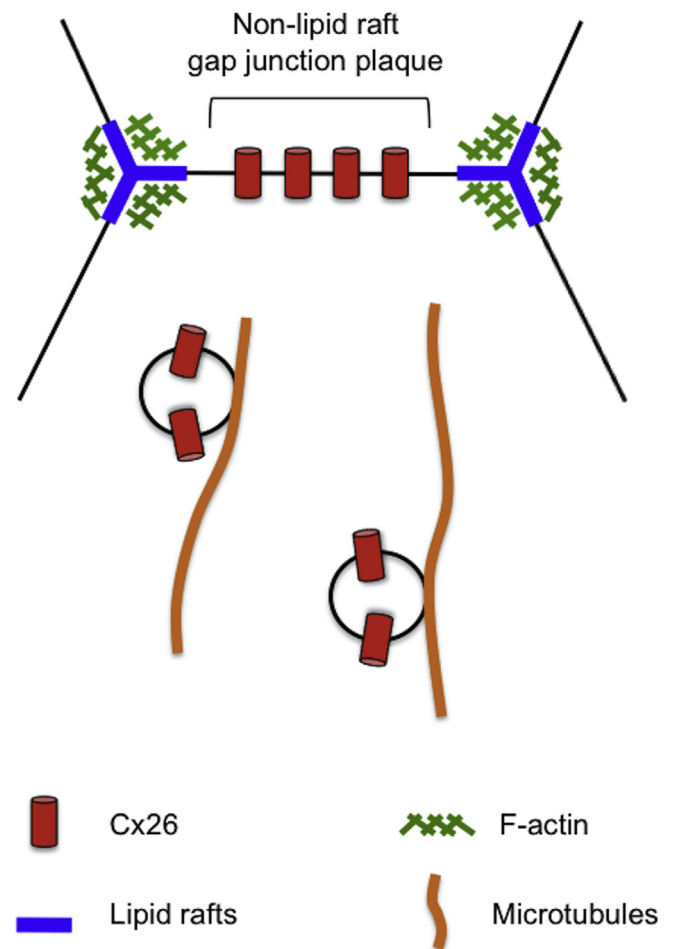
The composition of the plasma membrane lipid bilayer has a strong effect on ion channels (Cascio, 2005; Defamie and Mesnil, 2012). *In vitro* studies have shown that Cx26 has a low affinity for cholesterol and a preference for tight association with anionic phospholipids (Hung and Yarovsky, 2011; Locke and Harris, 2009). Lipid rafts, which are known to modulate gap junction communication (Defamie and Mesnil, 2012), contain a high quantity of cholesterol and a low amount of anionic phospholipids (Pike, 2003). Using Cholera Toxin B subunit as a relevant marker of these membrane microdomains (Janes et al., 1999), we tested whether Cx26 assembles into lipid rafts or not in inner sulcus cells. We found that Cx26 does not associate with lipid rafts, which are exclusively present at multicellular junction sites between adjacent GJPs (Fig. 2). Similar mutually exclusive localization patterns are observed in outer sulcus cells (Fig. S1B). These findings are consistent with previous data showing that Cx26 is excluded from lipid rafts in cell culture (Locke et al., 2005; Schubert et al., 2002). Moreover, we found that F-actin specifically associates with lipid rafts at multicellular junction sites, as expected (Chichili and Rodgers, 2009) (Fig. S3). Therefore, these data reveal an unusual submembrane compartmentalization for cochlear Cx26.

### 3.3. Actin-independent assembly of Cx26 into gap junction plaques

Distinct cytoskeletal tracks are required for trafficking lipid raft-associated or non-lipid raft-associated proteins to the plasma membrane (Jacob et al., 2003). Lipid raft-associated proteins travel to the cell periphery along microtubules and use actin filaments in the cell cortex to reach lipid raft membrane microdomains. In contrast, non-lipid raft-associated proteins traffic along microtubules and reach their plasma membrane destination without requiring actin tracks (Jacob et al., 2003). We then tested whether microtubules and/or actin networks are needed for delivery of Cx26 to non-lipid raft GJPs in inner sulcus cells. To this end, explants of organ of Corti isolated from newborn mice were treated with nocodazole and cytochalasin D as microtubules- and actin-disrupting drugs, respectively. Nocodazole treatment disturbed the transport of Cx26 to the cell periphery and strongly reduced the formation of GJPs in inner sulcus cells (Fig. 3B,D,E). In this condition, actin networks extend towards bicellular junctions, likely because microtubule depolymerization promotes reassembly of actin stress fibers (Ezratty et al., 2005). In agreement with previous *in vitro* data (George et al., 1999; Martin et al., 2001), an intact microtubule network is thus required for trafficking cochlear Cx26 from the interior of the cell to the plasma membrane. In contrast, we found that cytochalasin D did not significantly affect the assembly of Cx26 into GJPs (Fig. 3C–E). The effect of cytochalasin D on actin networks was ascertained by the presence of actin aggregates throughout the cytoplasm (Schliwa, 1982) (arrowhead on Fig. 3C). These findings suggest that, unlike other connexins, Cx26 uses an actin-independent trafficking pathway to assemble into non-lipid raft GJPs in the cochlea. This is consistent with recent data showing that Cx30 does not rely on actin network for GJP assembly when expressed as heteromeric channels with Cx26 in inner sulcus cells (Defourmy et al., 2019). In contrast, in Deiters' cells (i.e. the supporting cells which surround the outer hair cells), Cx30

homomeric channels associate with the actin network, which likely promotes the recruitment of Cx30 from the peripheral region to the GJP (Defourmy et al., 2019). Together, these results suggest that intrinsic Cx26 gap junction assembly features dominate the ones of Cx30 *in situ*.

Whereas most connexins assemble in a “two-step mechanism”, i.e. in a microtubule- and actin-dependent fashion, our data suggest that cochlear Cx26 most likely traffics in a kind of “one-step process”, only microtubule-dependent (Fig. 4). Such «actin bypass» pathway could be needed to ensure the relatively fast turnover rate of Cx26 (Traub et al., 1987). This could be the reason why Cx26 oligomers are already found in the endoplasmic reticulum (Diez et al., 1999) and can reach the cell surface via a route bypassing the Golgi complex (George et al., 1999; Martin et al., 2001). Although connexins usually reach the plasma membrane by traveling along the secretory pathway (Laird, 2006), this non-classical route used by Cx26 should be a faster process than trafficking through the Golgi (Grieve and Rabouille, 2011). In this context, an *in vitro* cell-free transcription/translation system has shown that Cx26 exhibits a singular membrane-integration behaviour and integrates directly in a post-translational manner into plasma membranes. Protein-cleavage studies of Cx26 integrated into plasma membranes indicated a similar native transmembrane topography to that of Cx26 integrated co-translationally



**Fig. 4. A model for how Cx26 traffics and assembles into GJPs in the cochlea.** Cx26 travels along microtubules and assembles into non-lipid raft GJPs in an actin-independent fashion. F-actin associates with lipid rafts at multicellular junction sites between adjacent GJPs.

into microsomes (Ahmad and Evans, 2002; Zhang et al., 1996). Cx26 oligomerization and assembly into hemichannels thus occurs independently of the conventional biogenesis of gap junctions involving connexin trafficking and oligomerization via membrane components of the secretory pathway (Ahmad and Evans, 2002).

Cx26 channel activity has been shown to be highly sensitive to cholesterol concentration (Locke and Harris, 2009). As a consequence, it should be of interest to determine how junctional Cx26 are specifically retained in non-lipid raft compartments. In a mouse embryo fibroblast cell line, the distribution of Cx43 at cell interfaces has been shown to be controlled by the transmembrane protein ephrin-B1 (Davy et al., 2006). Interestingly, a protein from the same family, ephrin-B2, was found to be specifically expressed in cochlear supporting cells (Defourny et al., 2015). Since gap junction communication is promoted at ephrin/ephrin interfaces within a cell compartment (Davy et al., 2006), it is tempting to speculate that ephrin-B2 might regulate Cx26 distribution and gap junction communication between adjacent supporting cells.

#### 4. Conclusion

Altogether, our data provide an unexpected insight into Cx26 trafficking pathway and gap junction assembly in the cochlea. Since many of the disease-causing mutations in *GJB2* impair the trafficking and delivery of Cx26 to the cell surface (Ambrosi et al., 2013; Hoang Dinh et al., 2009; Xu and Nicholson, 2013), our findings should help further research aimed to decipher the pathogenic significance of these mutations.

#### Competing financial interests

The authors declare no competing financial interests.

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#### Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.heares.2019.01.020>.

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